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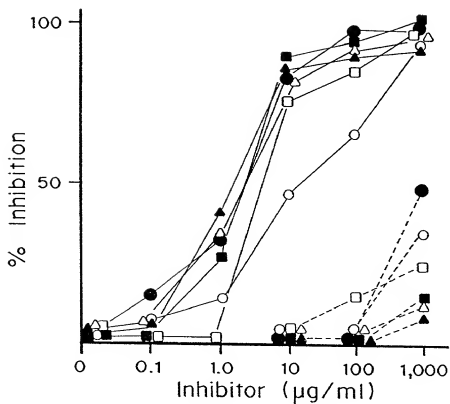


FIG. 1

FIG. 2a

GGT GAA GAA GGA GTT GTG CCA GCA CGT GAG TAC TCA GAC GAT CGT 45
 Gly Glu Glu Gly Val Val Pro Ala Arg Glu Tyr Ser Asp Asp Arg [15]

 AAC ATC AAC CTG GCA GAC GAA TTA AAA ATT GGT GAT ACC ATT GAA 90
 Asn Ile Asn Leu Ala Asp Glu Leu Lys Ile Gly Asp Thr Ile Glu [30]

 GCA GTT GTC ATT TCT AAC GTA ACA AGC GAC AAG GAA GGC GTC AGT 135
 Ala Val Val Ile Ser Asn Val Thr Ser Asp Lys Glu Gly Val Ser [45]

 TAC TTG TTG TCA AAG AAG CGT TTG GAT GCG CGC AAG GCA TGG GAA 180
 Tyr Leu Leu Ser Lys Lys Arg Leu Asp Ala Arg Lys Ala Trp Glu [60]

 AAC TTG AGC TTT GCT GAA GGT GAC ACA GTT GAT GCC AAG GTT ATC 225
 Asn Leu Ser Phe Ala Glu Gly Asp Thr Val Asp Ala Lys Val Ile [75]

 AAC GCT GTT CGT GGT GGT TTG ATT GTT GAT GTT AAC GGC GTA CGT 270
 Asn Ala Val Arg Gly Gly Leu Ile Val Asp Val Asn Gly Val Arg [90]

 GGT TTC GTA CCA GCA TCA ATG GTT GCA GAA CGT TTC GTT TCT GAT 315
 Gly Phe Val Pro Ala Ser Met Val Ala Glu Arg Phe Val Ser Asp [105]

 TTG AAC CAA TTC AAG AAT AAG GAT ATT AAA GCA CAA GTT ATC GAA 360
 Leu Asn Gln Phe Lys Asn Lys Asp Ile Lys Ala Gln Val Ile Glu [120]

 ATT GAC CCT GCT AAT GCA CGT TTG ATT TTG TCA CGT AAG GCT GTT 405
 Ile Asp Pro Ala Asn Ala Arg Leu Ile Leu Ser Arg Lys Ala Val [135]

 GCT GCA CAA GAA CGC GCT GCA CGA TTG GCT GAA GTA TTT AGC AAG 450
 Ala Ala Gln Glu Arg Ala Ala Gln Leu Ala Glu Val Phe Ser Lys [150]

 TTG TCA GTT CGT GAA GTT GTT GAA GGA ACT GTT GCC CGT TTG ACA 495
 Leu Ser Val Gly Glu Val Val Glu Gly Thr Val Ala Arg Leu Thr [165]

 GAC TTC GGC GCA TTC GTT GAC TTG GGT GGT GTT GAT GGT TTG GTT 540
 Asp Phe Gly Ala Phe Val Asp Leu Gly Gly Val Asp Gly Leu Val [180]

 CAC GTA TCA GAA ATC TCA CAC GAT CGT GTG AAG AAC CCG GCC GAT 585
 His Val Ser Glu Ile Ser His Asp Arg Val Lys Asn Pro Ala Asp [195]

 GTA TTG ACA AAG GGT GAC AAG GTT GAT GTT AAG ATC TTG GCA TTG 630
 Val Leu Thr Lys Gly Asp Lys Val Asp Val Lys Ile Leu Ala Leu [210]

 GAC ACT GAA AAG GGT CGT ATC TCA TTG TCA ATC AAA GCA ACA CAA 675
 Asp Thr Glu Lys Gly Arg Ile Ser Leu Ser Ile Lys Ala Thr Gln [225]

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[illegible]

FIG. 3a

HS1	63	SFAEGDTVDKVINAVRGGLIVDVGVRGFPVPSMVAERFVSDLNQFNKNDIKAQVIEI	121
ES1	101	AYEDAE--TGVMGK-K--FT-ELD-I-A-L-G-L-DV-P-R-TLHLEGELEFK--KL	159
RS1	107	K-EA-ER-EGIIF-Q-K--FT--LD-AVA-L-R-Q-DI-PIR-VTPADAQPAALRNILKM	165
PS1	1	E--TGVMGK-K--FT-EL-I-A-L-G-L-DV-P-R-TLHLEGELEFK--KL	54
CS1	182	--DVV-KG-IVG-NK--VWAL-E-L- ----F-QISSK-SAEELLE-E-PLKPV-V	236
		a b c	
		****	***
HS1	122	DPANARILSRKAVAAQERAAQLAEVFSKLSVGEWEGTVARLTDGAFVDLGGVDGLV	180
ES1	160	-QKRNNVWV--R-IES-NS-ERDQLEN-QE-ME-K-I-KN--Y-----L	218
RS1	166	-KRRGNWV--RT-LEES--E-RS-IVQN-EE-Q- --V-KNI-Y-----I--L	224
PS1	55	-QKRNNVWV--R-IES-SS-ERDQLEN-QE-ME-K-I-KN--Y-----L	113
CS1	237	-EEQS--VM-NRKAM-DSQ- M-DSQAQ-GI-S-T--QS-KPY--I-I--IN-L	287
		d e	
		****	****
HS1	181	HVSEISHDRVKNPADVLTKGDKVDVKIALDTEKGRISLSIKATQRGPWDEAADQIAAG	239
ES1	219	-ITDMAWK--H-SEINV--E-IT--V-KF-R-RT-V-GL-QLGED--VAI-KRYPE-	277
RS1	225	-TDMAWR--H-SEIQNI-QQ-K-Q-IRINQ-TH---GM-QLESD--GIGAKYPV-	283
PS1	114	-ITDMAWK--H-SEINV--E-IT--V-KF-R-RT-V-GL-QLGED--VAI-KRYPE-	172
CS1	288	---Q-----SDI-I--QP--TLK-M--SH-R-R-V--T-KLEPT-G-	337
		a b c d	
		****	****

FIG. 3b

HS1	240	SVLEGT	KRVKDF	GA	FEIL	P	IEGL	VHVS	QISN	KRIEN	PSEVL	KSGD	KVQV	KVLD	DIKP	298	
ES1	278	TK-T	-R-	TNLT	-Y-	C	----	EE	-V-	-----	EM	RD-V	-DATL	--SV	--E-EA	FTGVDR	510
RS1	284	KKIS	----	TNIT	-Y-	-----	LE	-----	I	-I-EM	-RPG	-QVI	-EFNK	--V-	RAV	--VDV	430
PS1	173	TK-T	-R-	TNLT	-Y-	C	----	EE	-V-	-----	EM						319

-K--EIAAV--QVDA

e

HS1	299	AERISL	SMKALE	EKP	PERE	317
ES1	511	KNPA	----	VR	-KD-AD	-KD 529
RS1	431	DK	----	GI	-QL	442
PS1	320	ER	----	GV	-QLA-DP	335

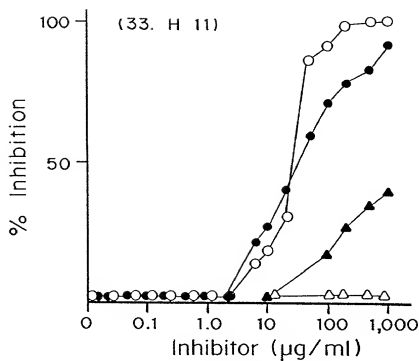


FIG. 4a

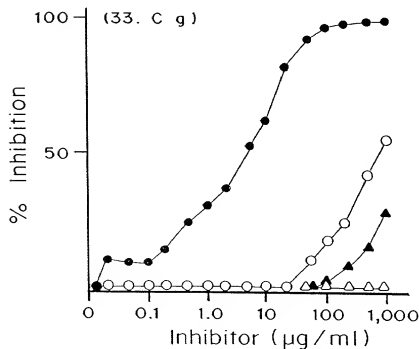


FIG. 4b

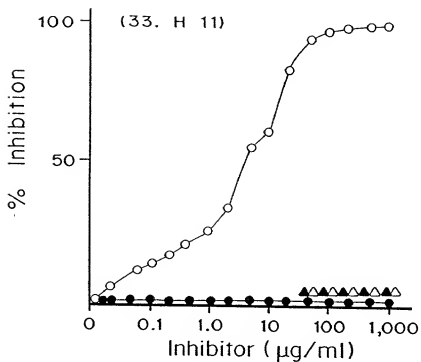


FIG. 4c

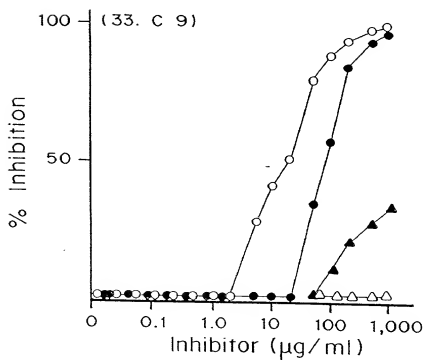


FIG. 4d